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Date 4/13/01 Label No. 02853599132US

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D B Peck
Name (Print)

D B Peck
Signature

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ACCOUNT NO. 04-0100

Docket No.:3380/11127-US4

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In Re Application of: Papsidero et al.

Serial No: TBA

Examiner: TBA

Filed: Concurrently Herewith

Group Art Unit: TBA

For: DETECTION AND TREATMENT OF BREAST DISEASE

**SUBMISSION OF SEQUENCE LISTING AND
STATEMENT PURSUANT TO 37 C.F.R. § 1.821**

Hon. Commissioner of Patents
Washington, DC 20231

Sir:

Pursuant to the requirements of 37 C.F.R. §1.821 through §1.825 for Sequence Listings, a computer readable form (diskette) containing the Sequence Listing and a paper copy of the Sequence Listing is enclosed herewith.

STATEMENT PURSUANT TO 37 C.F.R. § 1.821

Enclosed herewith is a computer readable form (diskette) containing sequence disclosure. Pursuant to Rule 37 C.F.R. §1.821, applicants herein state that

the contents of the labeled diskette, specifically the ASCII-encoded file therein labeled "Seqlist.txt", is identical to the paper copy of the sequence listing submitted herewith.

Respectfully submitted,

Dated: April 13, 2001


Neepa Y. Choksi
Registration No. 47,488
Agent for Applicant(s)

SEQUENCE LISTING

<110> Lawrence, Papsidero
Lyn, Dyster
Jana, Frustaci

<120> Detection and Treatment of Breast Cancer

<130> 3380/1I127-US4

<140> TBA
<141> Concurrently Herewith

<150> 09/146,580

<151> 1998-09-03

<150> 60/071,899

<151> 1998-01-20

<150> 60/092,155

<151> 1998-07-09

<160> 35

<170> PatentIn version 3.0

<210> 1

<211> 127

<212> PRT

<213> Homo sapiens

<220>

<221> UNSURE

<222> (70) .. (70)

<223> Xaa at position 70 is either Arg or Gly

<220>

<221> UNSURE

<222> (91) .. (91)

<223> Xaa at position 70 is either Lys or Asn

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Met Gln Gln Arg Gly Leu Ala Ile Val Ala Leu Ala Val Cys Ala Ala
1 5 10 15

Leu His Ala Ser Glu Ala Ile Leu Pro Ile Ala Ser Ser Cys Cys Thr
20 25 30

Glu Val Ser His His Ile Ser Arg Arg Leu Leu Glu Arg Val Asn Met
35 40 45

Cys Arg Ile Gln Arg Ala Asp Gly Asp Cys Asp Leu Ala Ala Val Ile
50 55 60

Leu His Val Lys Arg Xaa Arg Ile Cys Val Ser Pro His Asn His Thr
65 70 75 80

Val Lys Gln Trp Met Lys Val Gln Ala Ala Xaa Lys Asn Gly Lys Gly
85 90 95

Asn Val Cys His Arg Lys Lys His His Gly Lys Arg Asn Ser Asn Arg
100 105 110

Ala His Gln Gly Lys His Glu Thr Tyr Gly His Lys Thr Pro Tyr
115 120 125

<210> 2
<211> 104
<212> PRT
<213> Homo sapiens

<220>
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<222> (47) .. (47)
<223> Xaa at position 47 is either Arg or Gly

<220>
<221> UNSURE
<222> (68) .. (68)
<223> Xaa at position 47 is either Lys or Asn

<400> 2

Leu Pro Ile Ala Ser Ser Cys Cys Thr Glu Val Ser His His Ile Ser
1 5 10 15

Arg Arg Leu Leu Glu Arg Val Asn Met Cys Arg Ile Gln Arg Ala Asp
20 25 30

Gly Asp Cys Asp Leu Ala Ala Val Ile Leu His Val Lys Arg Xaa Arg
35 40 45

Ile Cys Val Ser Pro His Asn His Thr Val Lys Gln Trp Met Lys Val
50 55 60

Gln Ala Ala Xaa Lys Asn Gly Lys Gly Asn Val Cys His Arg Lys Lys

65 70 75 80
His His Gly Lys Arg Asn Ser Asn Arg Ala His Gln Gly Lys His Glu
85 90 95
Thr Tyr Gly His Lys Thr Pro Tyr
100

<210> 3
<211> 18
<212> PRT
<213> Homo sapiens

<400> 3

Thr Glu Val Ser His His Ile Ser Arg Arg Leu Leu Glu Arg Val Asn
1 5 10 15

Met Cys

<210> 4
<211> 16
<212> PRT
<213> Homo sapiens

<400> 4

Lys Asn Gly Lys Gly Asn Val Cys His Arg Lys Lys His His Gly Lys
1 5 10 15

<210> 5
<211> 19
<212> PRT
<213> Homo sapiens

<400> 5

Asn Ser Asn Arg Ala His Gln Gly Lys His Glu Thr Tyr Gly His Lys
1 5 10 15

Thr Pro Tyr

<210> 6
<211> 3117
<212> DNA
<213> Homo sapiens

<220>

<221> unsure
<222> (1)..(3117)
<223> n at any position in the sequence may represent a or g or c or t/

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agtgaatatg tgtcgcatcc agagagctga tggggattgt gacttggctg ctgtcatcct 240
tcatgtcaag cgcnngaagaa tctgtgtcag cccgcacaac catactgtta agcagtggat 300
gaaagtgcaa gctgccaana aaaatggtaa aggaaatgtt tgccacagga agaaaacacca 360
tggcaagagg aacagtaaca gggcacatca gggaaacac gaaacatacg gccataaaac 420
tccttattag agaatctaca gataaatcta cagagacaat ccccaagtg gacttggcca 480
tgattggttg taagtttatac atctgaattc tccttattgt agacaacaga acaaaacaaa 540
atattggttt ttaaaaaatg aacaattgtg ccgtatgcaa atgtacccaa taatatactc 600
cactggaaaa tgaatgaaa aaannatact ggctgggtat ggtgggtccc ccctttatc 660
ccannnnctt cgggaggcag aggcaggagg atcacttgag accaggantt ngagacnagc 720
tngggcaaa anagcaanga cntcattnt acaaacnaaa aaaaanntg gcccggcntg 780
gtagnacttgcntataatcc cagcnacatg ggaggtngag gtgggaggat cacttgagtc 840
tggngagtt ngaggtngca gtgagcagcn tgggtgacag aatgnagacc ntgtctctaa 900
aaataataat aataatgata gtgtatatct tcataataata ttttaagnag gagcatatag 960
atataacttn ctcccaactt tttaattata gtttccaaa cttacagaga agttaaaaga 1020
atggtacaat gaacatctat atatcttca ccacaatatt aatcattgtt aatattgtgc 1080
cacatttgct ttctctctcc tctcttgta ggggttncaa tataaaatata tataacttt 1140
aaaatataatc ttgttttgct aaccattgga aaataagttg caaaaatcat gacacttcac 1200
ccctagtttc tttnnggtgt tataacttga cataccctaa aataaagaca tttttctaca 1260
taatcacctt atcagttta tacctaaaaa attaataatt tcataataata tattccatat 1320

tcaaatttc ccaactattt agagaggcatt ttatgttagtt ttttttcac tccagtaatc 1380
aatcaaggtn gacatacata ttgcaaataa ttgttatttt tcttaatat cttaatct 1440
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tgctgtgtct tccaggctgg agtgcagtgg cacaatttg atttggctc actgaagcct 1560
caactttagg gctcaagcaa tcctcccacc tcagcctncc cgagtatctg ggatcaaggt 1620
gcataccac cacacctggc taatttgtt tattttgt agagacaggg tctcactatg 1680
ttgcccaggt tgatctcaa ctcctggct caagcgatcc tcccacctta gcctcccaa 1740
gtactggat tataggtgtg agccacagtg cctggctaa ttatTTctt gtgatcaa 1800
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agaacaaaaaa gcctgcctt ctgccccaa csnntgcyt cccccagtna gttcttgng 2040
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tgtttgtttt tttkgagaca grrtntnayt ctgttgccc argctggart tgcaagtggca 2220
caatynggy tncattgcaa cytngcytc cssgcgttc aaktgatyyt cttgcycatcag 2280
cytccccaaag taantgatat tacaggncc cagccaccam accccgntga wttttgtatt 2340
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ktgaaccacc cgcctgtgcc ycccaaantg ctggaattac cancgttgan ccaccatgcc 2460
gggcycacac gtttgarttt ganaccattg tnccattcct cttttggcct yttttttntc 2520
catagnngct tcaagataga tangtaagrg cccagtagtn gttcwtarga agcnmatagr 2580
rancrggarc canttnatc aggtggcag gtgtccnngg cytccctgct ggytnntccc 2640
aagcggtggt gttgccarga nktnttggar gtgataatgg gananaccag naggcmctga 2700
gtyncnntag gttnaaatgc cacaaaaact ggccttggc ctaatatccy ycmttgamta 2760
nttarcatc awtttattwa tttncctgac attntgcma ncctttgtwt ttntatttcc 2820

nctntatara	wgargaaatt	tgaggntytt	araggtaaaa	tganttgcnc	nrgtnnnacmc	2880
aggaagtggc	nraranaanc	tttttanatn	mgaaaaaaatt	aataaaatat	aatatgagag	2940
taacttaaaa	tattaataaa	ccacaatttt	aaattaatta	accgtgataa	ccaacattaa	3000
taaaagttaa	gataccaaaa	cactggtgtn	taatttttn	aactaacaan	ttgaattatt	3060
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<210> 7
<211> 381
<212> DNA
<213> Homo sapiens

<220>
<221> unsure
<222> (207)..(207)
<223> n may represent a or g or c or t/u

<220>
<221> unsure
<222> (272)..(272)
<223> n may represent a or g or c or t/u

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gaagccatac ttcccattgc ctccagctgt tgcacggagg tttcacatca tatttccaga 120
aggctcctgg aaagagtgaa tatgtgtcgc atccagagag ctgatggga ttgtgacttg 180
gctgctgtca tccttcatgt caagcgcnga agaatctgtc tcagcccgca caaccatact 240
gttaagcagt ggatgaaagt gcaagctgcc aaaaaaaatg gtaaaggaaa tgtttgccac 300
aggaagaaac accatggcaa gaggaacagt aacagggcac atcagggaa acacgaaaca 360
tacggccata aaactcctta t 381

<210> 8
<211> 104
<212> DNA
<213> Homo sapiens

<400> 8

acacgaattc acgttagaaaa ttcttaacca aaaacattaa acctgaattt gatcacaaga 60
aaataattag gccaggcact gtggctcaca cctataatcc cagt 104

<210> 9
<211> 25
<212> DNA
<213> Homo sapiens

<400> 9
gaattcacgt aggaaattct taacc 25

<210> 10
<211> 22
<212> DNA
<213> Homo sapiens

<400> 10
actgggattta taggtgtgag cc 22

<210> 11
<211> 311
<212> DNA
<213> Homo sapiens

<220>
<221> unsure
<222> (101)..(101)
<223> n may be a or g or c or t/u

<220>
<221> unsure
<222> (162)..(162)
<223> n may be a or g or c or t/u

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tccactgctt aacagtatgg ttgtgcgggc tgacacagat tnttctgcgc ttgacatgaa 180
ggatgacagc agccaagtca caatccccat cagctctctg gatgcgacac atattcactc 240
tttccaggag ccttctggaa atatgatgtg aaacacctcggt gcaacagctg gaggcaatgg 300

gaagtatggc t

311

<210> 12
<211> 20
<212> DNA
<213> Artificial

<220>
<223> Sequencing primer T7

<400> 12
taatacgact cactataggg

20

<210> 13
<211> 18
<212> DNA
<213> Artificial

<220>
<223> pCR3.1 Reverse Primer

<400> 13
tagaaggcac agtcgagg

18

<210> 14
<211> 22
<212> DNA
<213> Artificial

<220>
<223> Gene specific primer (24R)

<400> 14
actgggattta taggtgtgag cc

22

<210> 15
<211> 24
<212> DNA
<213> Artificial

<220>
<223> Gene specific primer (24R2)

<400> 15
caaattcagg tttaatgttt ttgg

24

<210> 16
<211> 21
<212> DNA
<213> Artificial

<220>
<223> Gene specific primer (F4)

<400> 16
ctcaaacgtg tgagccggc a

21

<210> 17
<211> 25
<212> DNA
<213> Artificial

<220>
<223> Gene specific primer (F3)

<400> 17
gtactcaaa ctagacgttt tgaag

25

<210> 18
<211> 24
<212> DNA
<213> Artificial

<220>
<223> primers F8

<400> 18
ccgtatgttt cgtgtttccc ctga

24

<210> 19
<211> 24
<212> DNA
<213> Artificial

<220>
<223> Primer R5

<400> 19
agccatactt cccattgcct ccag

24

<210> 20
<211> 150

<212> PRT

<213> Homo sapiens

<400> 20

Met Asn Leu Trp Leu Leu Ala Cys Leu Val Ala Gly Phe Leu Gly Ala
1 5 10 15

Trp Ala Pro Ala Val His Thr Gln Gly Val Phe Glu Asp Cys Cys Leu
20 25 30

Ala Tyr His Tyr Pro Ile Gly Trp Ala Val Leu Arg Arg Ala Trp Thr
35 40 45

Tyr Arg Ile Gln Glu Val Ser Gly Ser Cys Asn Leu Pro Ala Ala Ile
50 55 60

Phe Tyr Leu Pro Lys Arg His Arg Lys Val Cys Gly Asn Pro Lys Ser
65 70 75 80

Arg Glu Val Gln Arg Ala Met Lys Leu Leu Asp Ala Arg Asn Lys Val
85 90 95

Phe Ala Lys Leu His His Asn Met Gln Thr Phe Gln Ala Gly Pro His
100 105 110

Ala Val Lys Lys Leu Ser Ser Gly Asn Ser Lys Leu Ser Ser Ser Lys
115 120 125

Phe Ser Asn Pro Ile Ser Ser Lys Arg Asn Val Ser Leu Leu Ile
130 135 140

Ser Ala Asn Ser Gly Leu
145 150

<210> 21

<211> 95

<212> PRT

<213> Homo sapiens

<400> 21

Met Cys Cys Thr Lys Ser Leu Leu Leu Ala Ala Leu Met Ser Val Leu
1 5 10 15

Leu Leu His Leu Cys Gly Glu Ser Glu Ala Ser Asn Phe Asp Cys Cys
20 25 30

Leu Gly Tyr Thr Asp Arg Ile Leu His Pro Lys Phe Ile Val Gly Phe
35 40 45

Thr Arg Gln Leu Ala Asn Glu Gly Cys Asp Ile Asn Ala Ile Ile Phe
50 55 60

His Thr Lys Lys Lys Leu Ser Val Cys Ala Asn Pro Lys Gln Thr Trp
65 70 75 80

Val Lys Tyr Ile Val Arg Leu Leu Ser Lys Lys Val Lys Asn Met
85 90 95

<210> 22

<211> 94

<212> PRT

<213> Homo sapiens

<400> 22

Met Ala Pro Leu Lys Met Leu Ala Leu Val Thr Leu Leu Leu Gly Ala
1 5 10 15

Ser Leu Gln His Ile His Ala Ala Arg Gly Thr Asn Val Gly Arg Glu
20 25 30

Cys Cys Leu Glu Tyr Phe Lys Gly Ala Ile Pro Leu Arg Lys Leu Lys
35 40 45

Thr Trp Tyr Gln Thr Ser Glu Asp Cys Ser Arg Asp Ala Ile Val Phe
50 55 60

Val Thr Val Gln Gly Arg Ala Ile Cys Ser Asp Pro Asn Asn Gln Arg
65 70 75 80

Val Lys Asn Ala Val Lys Tyr Leu Gln Ser Leu Glu Arg Ser
85 90

<210> 23

<211> 96

<212> PRT

<213> Homo sapiens

<400> 23

Met Gln Ile Ile Thr Thr Ala Leu Val Cys Leu Leu Leu Ala Gly Met
1 5 10 15

Trp Pro Glu Asp Val Asp Ser Lys Ser Met Gln Val Pro Phe Ser Arg
20 25 30

Cys Cys Phe Ser Phe Ala Glu Gln Glu Ile Pro Leu Arg Ala Ile Leu
35 40 45

Cys Tyr Arg Asn Thr Ser Ser Ile Cys Ser Asn Glu Gly Leu Ile Phe
50 55 60

Lys Leu Lys Arg Gly Lys Glu Ala Cys Ala Leu Asp Thr Val Gly Trp
65 70 75 80

Val Gln Arg His Arg Lys Met Leu Arg His Cys Pro Ser Lys Arg Lys
85 90 95

<210> 24

<211> 77

<212> PRT

<213> Homo sapiens

<400> 24

Ala Gln Pro Asp Ser Val Ser Ile Pro Ile Thr Cys Cys Phe Asn Val
1 5 10 15

Ile Asn Arg Lys Ile Pro Ile Gln Arg Leu Glu Ser Tyr Thr Arg Ile
20 25 30

Thr Asn Ile Gln Cys Pro Lys Glu Ala Val Ile Phe Lys Thr Lys Arg
35 40 45

Gly Lys Glu Val Cys Ala Asp Pro Lys Glu Arg Trp Val Arg Asp Ser
50 55 60

Met Lys His Leu Asp Gln Ile Phe Gln Asn Leu Lys Pro
65 70 75

<210> 25

<211> 98

<212> PRT

<213> Homo sapiens

<400> 25

Met Lys Val Ser Ala Val Leu Leu Cys Leu Leu Leu Met Thr Ala Ala
1 5 10 15

Phe Asn Pro Gln Gly Leu Ala Gln Pro Asp Ala Leu Asn Val Pro Ser
20 25 30

Thr Cys Cys Phe Thr Phe Ser Ser Lys Lys Ile Ser Leu Gln Arg Leu
35 40 45

Lys Ser Tyr Val Ile Thr Thr Ser Arg Cys Pro Gln Lys Ala Val Ile
50 55 60

Phe Arg Thr Lys Leu Gly Lys Glu Ile Cys Ala Asp Pro Lys Glu Lys
65 70 75 80

Trp Val Gln Asn Tyr Met Lys His Leu Gly Arg Lys Ala His Thr Leu
85 90 95

Lys Thr

<210> 26

<211> 97

<212> PRT

<213> Homo sapiens

<400> 26

Met Lys Val Ser Ala Ala Leu Leu Trp Leu Leu Leu Ile Ala Ala Ala
1 5 10 15

Phe Ser Pro Gln Gly Leu Ala Gly Pro Ala Ser Val Pro Thr Thr Cys
20 25 30

Cys Phe Asn Leu Ala Asn Arg Lys Ile Pro Leu Gln Arg Leu Glu Ser
35 40 45

Tyr Arg Arg Ile Thr Ser Gly Lys Cys Pro Gln Lys Ala Val Ile Phe
50 55 60

Lys Thr Lys Leu Ala Lys Asp Ile Cys Ala Asp Pro Lys Lys Lys Trp
65 70 75 80

Val Gln Asp Ser Met Lys Tyr Leu Asp Gln Lys Ser Pro Thr Pro Lys
85 90 95

Pro

<210> 27

<211> 99

<212> PRT

<213> Homo sapiens

<400> 27

Met Lys Ala Ser Ala Ala Leu Leu Cys Leu Leu Leu Thr Ala Ala Ala
1 5 10 15

Phe Ser Pro Gln Gly Leu Ala Gln Pro Val Gly Ile Asn Thr Ser Thr
20 25 30

Thr Cys Cys Tyr Arg Phe Ile Asn Lys Lys Ile Pro Lys Gln Arg Leu
35 40 45

Glu Ser Tyr Arg Arg Thr Thr Ser Ser His Cys Pro Arg Glu Ala Val
50 55 60

Ile Phe Lys Thr Lys Leu Asp Lys Glu Asp Cys Ala Asp Pro Thr Gln
65 70 75 80

Lys Trp Val Gln Asp Pro Met Lys His Leu Asp Lys Lys Thr Gln Thr
85 90 95

Pro Lys Leu

<210> 28

<211> 99

<212> PRT

<213> Homo sapiens

<400> 28

Met Lys Val Ser Ala Ala Leu Leu Cys Leu Leu Leu Thr Ala Ala Ala
1 5 10 15

Phe Ile Pro Gln Gly Leu Ala Gln Pro Asp Ala Ile Asn Ala Pro Val
20 25 30

Thr Cys Cys Tyr Asn Phe Thr Asn Arg Lys Ile Ser Val Gln Arg Leu
35 40 45

Ala Ser Tyr Arg Arg Ile Thr Ser Ser Lys Cys Pro Lys Glu Ala Val
50 55 60

Ile Phe Lys Thr Ile Val Ala Lys Glu Asp Cys Ala Asp Pro Lys Gln
65 70 75 80

Lys Trp Val Gln Asp Ser Met Asp His Leu Asp Lys Gln Thr Gln Thr
85 90 95

Pro Lys Thr

<210> 29

<211> 91

<212> PRT

<213> Homo sapiens

<400> 29

Met Lys Val Ser Ala Ala Arg Leu Ala Val Ile Leu Ile Ala Thr Ala
1 5 10 15

Leu Cys Ala Pro Ala Ser Ala Ser Pro Tyr Ser Ser Asp Thr Thr Pro
20 25 30

Cys Cys Phe Ala Tyr Ile Ala Arg Pro Leu Pro Arg Ala His Ile Lys
35 40 45

Glu Tyr Phe Tyr Thr Ser Gly Lys Cys Ser Asn Pro Ala Val Val Phe
50 55 60

Val Thr Arg Lys Asn Arg Gln Val Cys Ala Asn Pro Glu Lys Lys Trp
65 70 75 80

Val Arg Glu Tyr Ile Asn Ser Leu Glu Met Ser
85 90

<210> 30
<211> 93
<212> PRT
<213> Homo sapiens

<400> 30

Met Lys Ile Ser Val Ala Ala Ile Pro Phe Phe Leu Leu Ile Thr Ile
1 5 10 15

Ala Leu Gly Thr Lys Thr Glu Ser Ser Ser Arg Gly Pro Tyr His Pro
20 25 30

Ser Glu Cys Cys Phe Thr Tyr Thr Tyr Lys Ile Pro Arg Gln Arg
35 40 45

Ile Met Asp Tyr Tyr Glu Thr Asn Ser Gln Cys Ser Lys Pro Gly Ile
50 55 60

Val Phe Ile Thr Lys Arg Gly His Ser Val Cys Thr Asn Pro Ser Asp
65 70 75 80

Lys Trp Val Gln Asp Tyr Ile Lys Asp Met Lys Glu Asn
85 90

<210> 31
<211> 92
<212> PRT
<213> Homo sapiens

<400> 31

Met Lys Leu Cys Val Thr Val Leu Ser Leu Leu Met Leu Val Ala Ala
1 5 10 15

Phe Cys Ser Pro Ala Leu Ser Ala Pro Met Gly Ser Asp Pro Pro Thr
20 25 30

Ala Cys Cys Phe Ser Tyr Thr Ala Arg Lys Leu Pro Arg Asn Phe Val
35 40 45

Val Asp Tyr Tyr Glu Thr Ser Ser Leu Cys Ser Gln Pro Ala Val Val
50 55 60

Phe Gln Thr Lys Arg Ser Lys Gln Val Cys Ala Asp Pro Ser Glu Ser
65 70 75 80

Trp Val Gln Glu Tyr Val Tyr Asp Leu Glu Leu Asn
85 90

<210> 32

<211> 93

<212> PRT

<213> Homo sapiens

<400> 32

Met Gln Val Ser Thr Ala Ala Leu Ala Val Leu Leu Cys Thr Met Ala
1 5 10 15

Leu Cys Asn Gln Val Leu Ser Ala Pro Leu Ala Ala Asp Thr Pro Thr
20 25 30

Ala Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe Ile
35 40 45

Ala Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Ser Val Ile
50 55 60

Phe Leu Thr Lys Arg Gly Arg Gln Val Cys Ala Asp Pro Ser Glu Glu
65 70 75 80

Trp Val Gln Lys Tyr Val Ser Asp Leu Glu Leu Ser Ala
85 90

<210> 33

<211> 92

<212> PRT

<213> Homo sapiens

<400> 33

Met Gln Val Ser Thr Ala Ala Leu Ala Val Leu Leu Cys Thr Met Ala
1 5 10 15

Leu Cys Asn Gln Phe Ser Ala Ser Leu Ala Ala Asp Thr Pro Thr Ala
20 25 30

Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe Ile Ala
35 40 45

Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Gly Val Ile Phe
50 55 60

Leu Thr Lys Arg Ser Arg Gln Val Cys Ala Asp Pro Ser Glu Glu Trp
65 70 75 80

Val Gln Lys Tyr Val Ser Asp Leu Glu Leu Ser Ala
85 90

<210> 34

<211> 89

<212> PRT

<213> Homo sapiens

<400> 34

Met Lys Gly Leu Ala Ala Ala Leu Leu Val Leu Val Cys Thr Met Ala
1 5 10 15

Leu Cys Ser Cys Ala Gln Val Gly Thr Asn Lys Glu Leu Cys Cys Leu
20 25 30

Val Tyr Thr Ser Trp Gln Ile Pro Gln Lys Phe Ile Val Asp Tyr Ser
35 40 45

Glu Thr Ser Pro Gln Cys Pro Lys Pro Gly Val Ile Leu Leu Thr Lys
50 55 60

Arg Gly Arg Gln Asp Cys Ala Asp Pro Asn Lys Lys Trp Val Gln Lys
65 70 75 80

Tyr Ile Ser Asp Leu Lys Leu Asn Ala
85

<210> 35

<211> 104

<212> DNA

<213> Homo sapiens

<400> 35

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60

aaataattag gccaggcact gtggctcaca cctataatcc cagt

104